

Amendments to the Specification:

Please replace the paragraph beginning at page 6, line 15, with the following rewritten paragraph:

-- Particularly conserved regions and amino acid residues common to nectin polypeptides were identified by aligning nectin polypeptide sequences with each other and additional closely-related members of the nectin-Ig superfamily of proteins. The amino acid sequence of nectin-3 α and nectin-4 (SEQ ID Nos: 6 and 24) were compared with the amino acid sequences of other nectin and Ig family members (SEQ ID NO:20, 22, and 25), using a multiple sequence alignment program. The alignment of these sequences is shown in Table 2, and includes consensus residues (capitalized), which are identical among at least a majority (three) of the five amino acid sequences in the alignment. In addition, lower case residues are shown on a separate line of Table 2 and represent residues that are not consensus residues, but are identical between human nectin-3 α and human nectin-4 (SEQ ID Nos: 6 and 24). --

Please replace Table 2 beginning at page 7, line 7, with the following rewritten Table:

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Table 2
Conserved Nectin Amino Acids

(Hs=Homo sapiens)
(Mus=Murine)

Protein (SEQ ID NO)	
HUNECTIN2 ([SEQ ID NO:122]) HUCD155 ([SEQ ID NO:125]) HUNECTIN1 ([SEQ ID NO:120]) HUNECTIN3 ([SEQ ID NO:16]) HUNECTIN4 ([SEQ ID NO:124]) consensus	<div> <div> MARAAALLPS RSPPTPLLWP LLLLLLL... MARAMAAAWP LLLVALLVLS L...ALGLTA LQPPPTPPPLL LLLFPLLLFS AEMWGPEAWL LLLLLLASFT </div> <div> P W </div> <div> 1 1 f </div> </div>
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	<div> <div> 51 100 </div> <div> ..ETGAQDVR VQVLPEVRGQ LGGTVELPCH L.LPPVPGLY ISLVTWQRPD WPPPGTGDDV VQAPTQVPGF LGDSVTLPCY LQVPNMEVTH VSQLTWAR.. FFLPGVHSQV VQVNDSEMYGF IGTDVVLHCS FANP.LPSVK ITQVTWQK.S RLCGALAGP. IIVEPHVTAV WGKNVSLKCL I..EV..NET ITQISWEKIH GRCP..AGE. LETSDVVTVV LGQDAKLPCF YRGDS..GEQ VGQVAWARVD [cPG ag VQV VtGv LG V LPC P e I QV W R] PG VQV V G LG V LPC P I QVTW R c ag t v e </div> </div>
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	<div> <div> 101 150 </div> <div> APANHQNVA FHPKMGPSFP SPKPGSERLS FVSAKQSTGQ DTEAELQDAT .HGESGSMV FHQTQGPSYS E...SKRLE FVAARLG...AELRNAS TNGSKQNVAI YNPSMGVSV. .LAPYRERVE FL.....RPSFTDGT .GKSSQTVAV HHPQYGFVSQ ..GEYQGRVL FKNYSLN...DAT AGEGAQELAL LHSKYGLHVS ..PAYEGRVE QPPPPRNPL.DGS [g Q A H yG SV Y gRVE F n DAT] Q VA HP G SV Y RVE F DAT g v g n </div> </div>

HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	151 LALHGLTVED EGNITCEFAT FPKGSVRGMT WLRVIAKPKN QAEAQKVTF. LRMFGLRVED EGNITCLFVT FPQGSRSVDI WLRVLAKPQN TAEVQKVQL. IRLSRLELED EGVYICEFAT FPTGNRESQL NLTVMKAPT N WIEGTQAVLR ITLHNIGFSD SGKYICKAVT FPLGNAQSST TVTVLVEPTV SLIKGPDSLI VLLRNAVQAD EGEYECRVST FPAGSFQARL RLRVLVPLP SLNPGP.ALE [L nL ED EG Y C F T FP GS q LRVLAKP N s E L] L L ED EG Y C F T FP GS LRVLAKP N E L n q v sl gp	200
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	201SQDPTT VALCISKEGR PPARISWLSS LDWEAKETQV SGTLAGTVTVTGEVPV MARCVSTGGR PPAQITWHS LGGMPNTSQV PGFLSGTVTV AKKGQDDKVL VATCTSANGK PPSVSVSWETR LKGEARVPGD SGTPMAPVTV DGGNE...TV AAICIAATGK PVAHIDWEGD LGEM..ESTT TSFPNETATI EGQGL...TL AASC.TAEGS PAPSVTWDTE VKGT..TSSR SFKHSRSAAV [g T aa C Sa G PPA I W L G S SG TVTV] T A C SA G PPA I W L G S SG TVTV g a a	250
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	251 TSRFTLVPSG RADGVTVTCK VEH..ESFEE PALIPVTLVS RYPPEVSISG TSLWILVPSS QVDGKNVTCK VEH..ESFEK PQLLTVNLTV YYPEVSISG ISRYRLVPSR EAHQQSLACI VNYHMDRFKESLTLNV QYEPEVTIEG ISQYKLPFTR FARGRRITCV VKHP..ALEK DIRYSFILDI QYAPEVSVTG TSEFHLVPSR SMNGQLTTCV VSHP..GLLQ DQRITHILHV SFLAEASVRG [TS LVPSR A G TC V Hp FE d r iL V Y PEVSI G] TS LVPSR A G TC V H FE L V Y PEVSI G v p l d r i v	300
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	301 Y.DDN.WYLG RTDATLSCDV RSNPEPTGYD WSTTSGTFPT SAVAQGSQLV Y.DDN.WYLG QNEATLTCD A RSNPEPTGYN WSTTMGPLPP FAVAQGAQLL F.DGN.WYLG RMDVKLTCKA DANPPATEYH WTTLNGSLPK GVEAQNRTLF Y.DGN.WFVG RKGVNLCNA DANPPPFKSV WSRLDGQWPD GLLASDNTLH LEDQNLWHIG REGAMLKCLS EGQPPPSYN. WTRLDGPLPS GVRVDGDTLG [YD N WYLG R gA LkC A NPPPTY WSTLdG LP G AQG TL] Y D N WYLG R A L C A NPPPT Y WSTL G LP G AQG TL g k r d	350
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	351 IH.AVDSLFFN TTFVCTVTNA VGMGRAEQVI FVRETP.... IR.PVDKPIN TTLICNVTA LGARQAELTV QVKEGP.... FKGPINYSLA GTYICEATNP IGTRSGQVEV NITEFPYTPS FVHPLTFNYS GVIYICKVTNS LGQRSDQKVI YISDPPTTT LQPTIQWHPS F.PPLTTEHS GIYVCHVSNE FSSRDSQVTV DVLDPQEDSG KQ..... [F Plt s G YIC VTN G R Q V EpP q] F P GTYIC VTN G R Q V V E P lt s dp g	400
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	401RAS P...RDV..G PLVWGA VGGT LLVLLLLLAGGPSE H...SGISRN AIIFLVLG... ILVFLILLGIPPE HGRRAGPVPT AIIGGVAGSI LLVLIVVGGI TADIEDLATE PKKLFPFLST LATIKDDTIA TIIASVVGGA LFIVLVSVLAVDLV...SAS VVVVGVI AAL LFCLLVVVV [d II GV G LLVLLV vG] dl f v	450
HUNECTION2 HUCD155 HUNECTION1 HUNECTION3 HUNECTION4	451 SLAFILLRVR RR....RKS .PGGAGGGAS GDGGFYDPKA QVLGNGDPVF GIYFYWSKCS REVLWHCHLC .PSSEHHQSC RN~~~~~ VVALRRRRRT FKGDYSTKKH .VYGNNGYSKA GIPQHHPMA QNLQYPPDS GIFCYRRRRRT FRGDYFAKNY IPPSDMQKES QIDVLQODEL D..SYP.DSV LMSRYHRR.. .KAQQMTQKY EEELTLTREN SIRRLHSHHT DPRSQPEESV [y RR y P e I lH d Ls PD Sv] Y RR P I P S y e l d s v	500

HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	501 WTPVVPGPME P.DGKDEEEE EEEKAEKGL MLPPPPALED DMESQLDGSL ~~~~~ .DEKKAGPLG G.SSYEEEE EEEGGGGGER KVGGPHPKYD EDAKRPYFTV .KKENKNP..VNNLIRKDY LEEPEKTOWN NVENLNRFER PMDYEDLKM GLRAEGHPDS LKDNSSCSVM SEEPEGRSYS TLTTVREIET QT...ELLSP [p n Eepe e 1] P EE E n pe e 1	550
HUNECTIN2 HUCD155 HUNECTIN1 HUNECTIN3 HUNECTIN4	551 ISRRVYV~ ~~~~~ ~~~~~ DEAEARQDGY GDRTLGYQYD PEQLDLAENM VSQNDGSFIS KKEWYV~ GM.KFVSDEH YDENEDDLVS HV...DGSVI SR...REWYV ~~~~~ GSGRAEEEEED QDEGIQAMN HFVQENGTLR AKPTGNGIYI NGRGHLV [g A E DE I H g y] A D g e e h g y	597